

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING**

### **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: \_\_\_\_\_

Source: \_\_\_\_\_

Date Processed by STIC: \_\_\_\_\_

10/613, 736A  
JFW/16  
08/01/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>**> , **EFS Submission User Manual** - ePAVE)
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U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



## RAW SEQUENCE LISTING

DATE: 08/01/2006

PATENT APPLICATION: US/10/613,736A

TIME: 13:53:28

Input Set : A:\C1037.70044US00.txt; Revised SEQ listing.txt

Output Set: N:\CRF4\08012006\J613736A.raw

375 &lt;220&gt; FEATURE:

376 &lt;221&gt; NAME/KEY: misc\_feature

377 &lt;222&gt; LOCATION: (27)..(46)

378 &lt;223&gt; OTHER INFORMATION: n is a, c, g, or t; and any 0-20 may be absent

381 &lt;400&gt; SEQUENCE: 26

E--&gt; 382 gggnnnnnnnn nnnnnnnnnnn nnnngggnnnnn nnnnnnnnnnn nnnnnnnnggg

49

10/6/13, 736 A

pg-3

<210> 24  
<211> 10  
<212> DNA  
<213> Artificial sequence

<220>

<223> Oligodeoxynucleotide

<400> 24  
tttttttcga

10

13

P/s delete

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/613,736A

DATE: 08/01/2006  
TIME: 13:53:29

Input Set : A:\C1037.70044US00.txt; Revised SEQ listing.txt  
Output Set: N:\CRF4\08012006\J613736A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:24; Line(s) 338

**VERIFICATION SUMMARY**

DATE: 08/01/2006

PATENT APPLICATION: US/10/613,736A

TIME: 13:53:29

Input Set : A:\C1037.70044US00.txt; Revised SEQ listing.txt

Output Set: N:\CRF4\08012006\J613736A.raw

L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
L:191 M:283 W: Missing Blank Line separator, <220> field identifier  
L:338 M:254 E: No. of Bases conflict, LENGTH:Input:13 Counted:10 SEQ:24  
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
L:357 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:26 SEQ:25  
L:382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0  
L:382 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:49 SEQ:26